

EFFECT OF STRONG N-TH COUPLING ON CORE DESIGN CALCULATIONS BASED ON A TYPICAL 100 MWE INTEGRAL PWR DESIGN

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Nuclear science and thechnology
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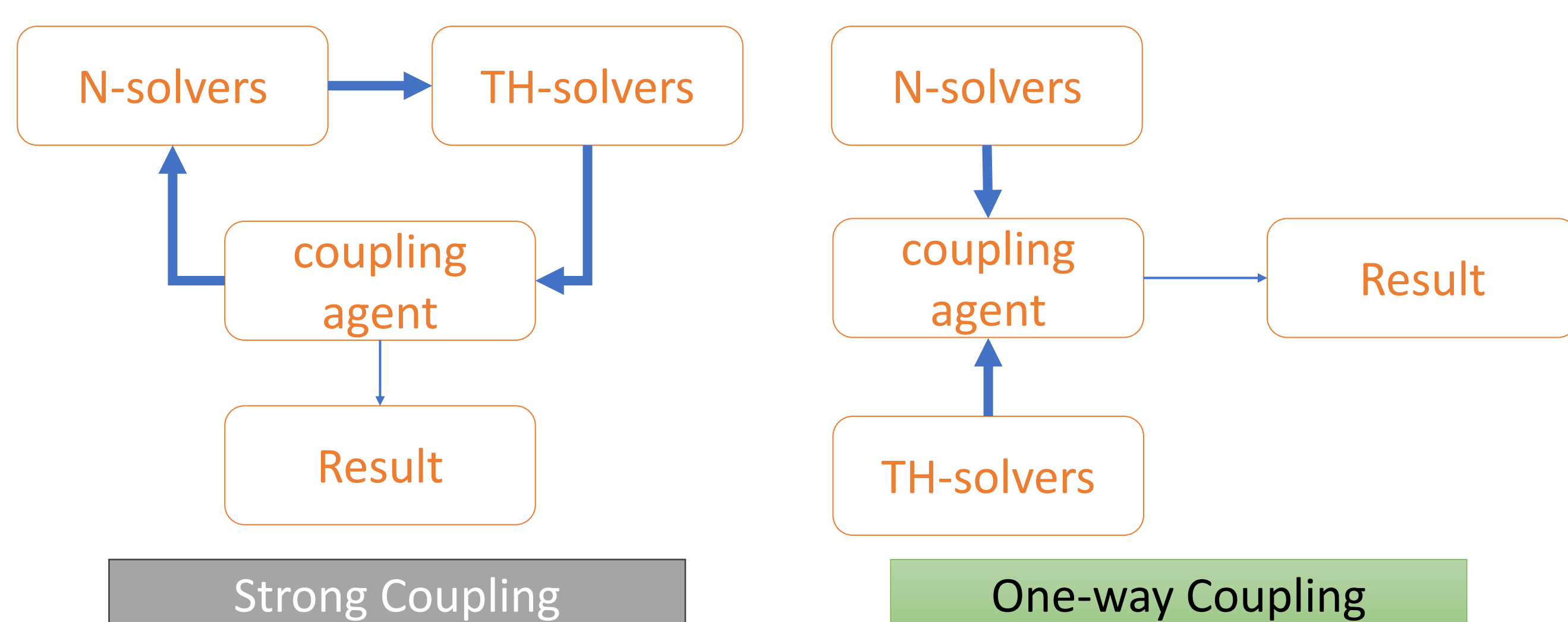
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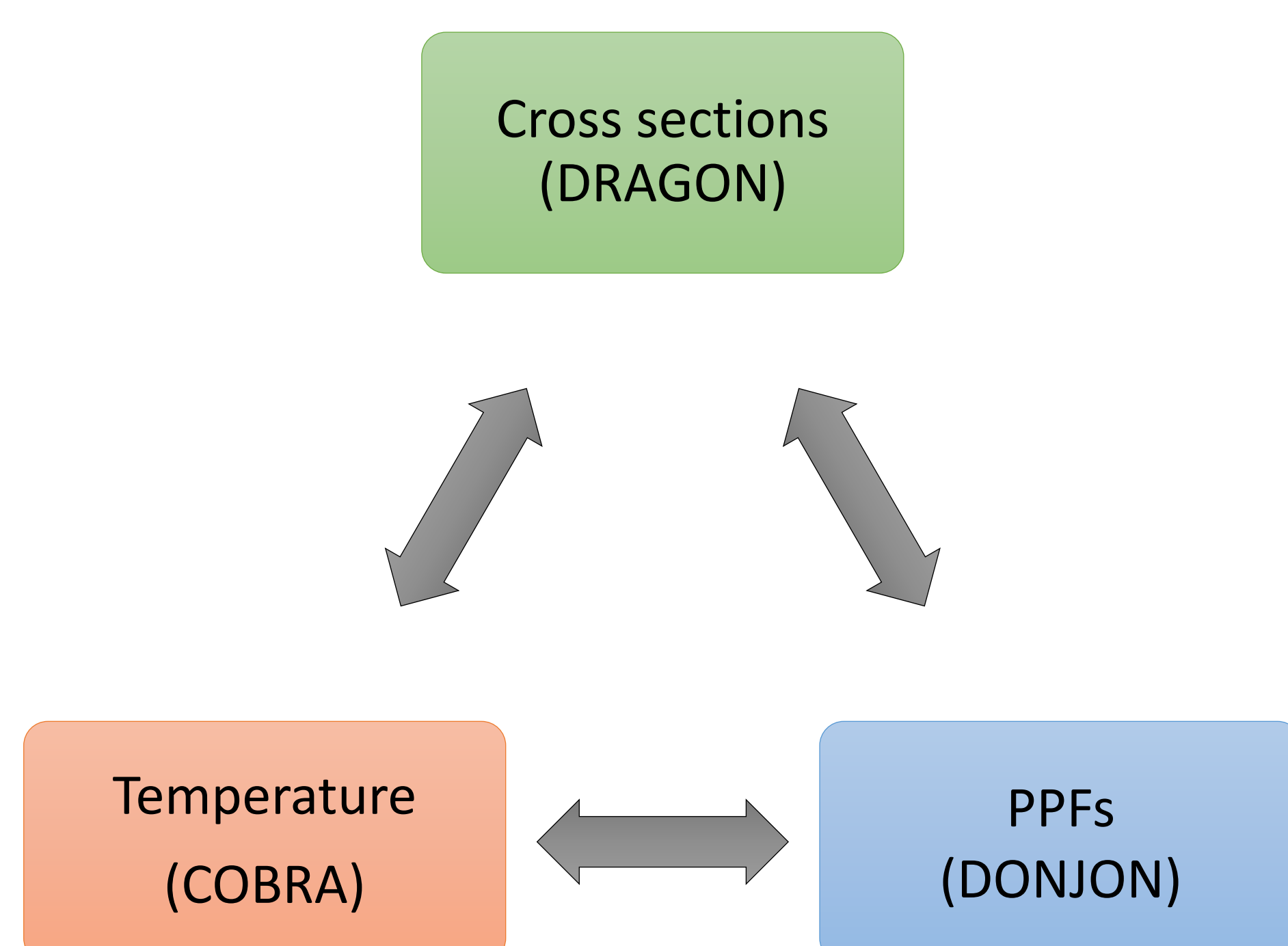
INTRODUCTION

- Reactor core **design** relies on simulation techniques
- Neutronics and thermal-hydraulics software tools are used
- Most of these software tools (codes) are **interdependent**
- N-calculations rely on the nuclear cross section data
- TH-calculations for safety parameters such as CHF, MDNBR
- One-way coupling: N and TH solvers are utilized independently.
- Strong coupling: N and TH solvers are interconnected.
- Aim: Comparison between **one-way** vs strong coupling
- Aim: the core of a typical 100 MWe SMR



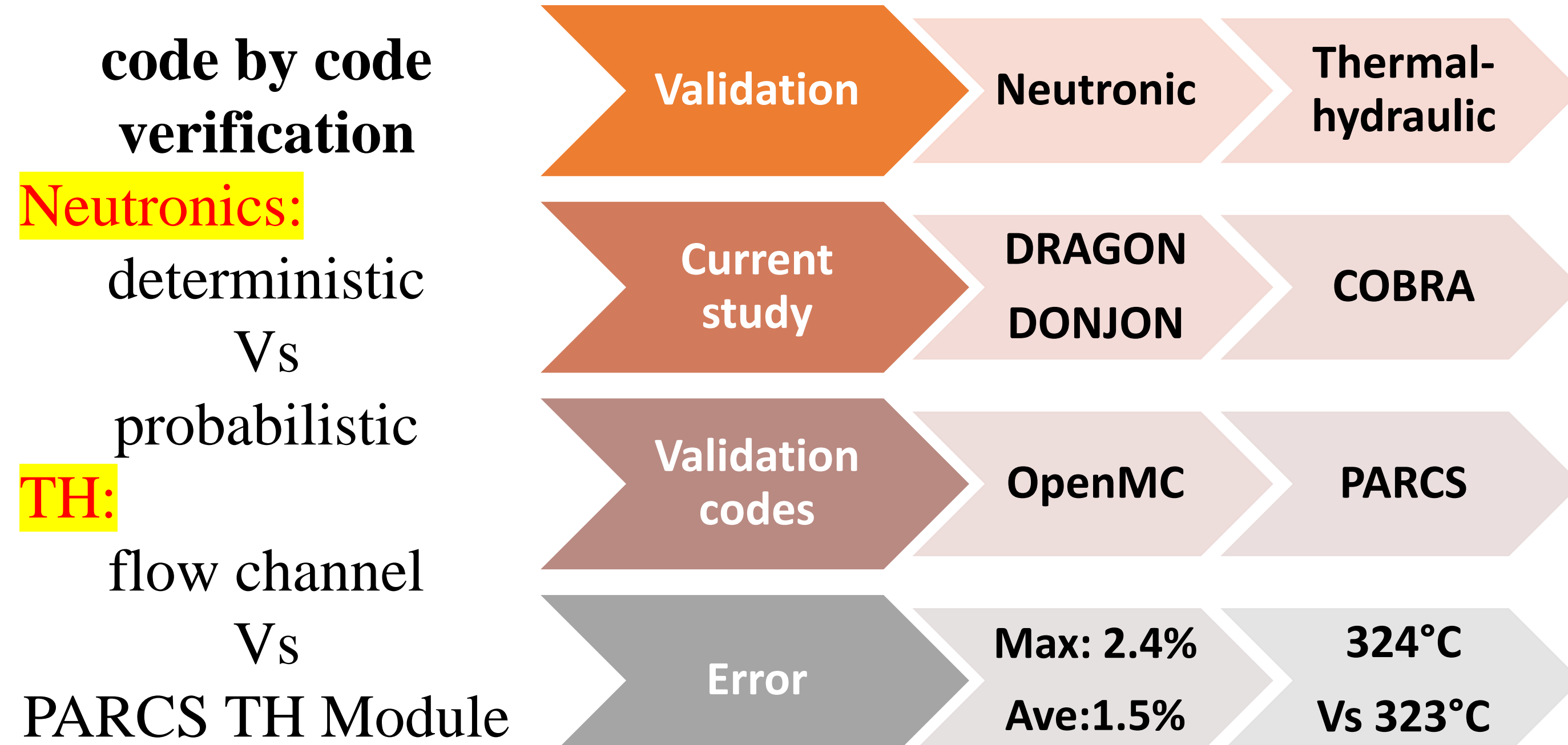
METHODS AND MATERIALS

- Neutronic cell calculations (DRAGON): cross sections
- Neutronic core calculations (DONJON): PPFs
- Thermal-hydraulic calculations (COBRA): Temperature



- The convergence is checked
- Above stages are iterated to reach convergence criteria

RESULTS



K_{eff} at the BOC

- Simple coupling \rightarrow 1.000593
- SAR \rightarrow 1.005724
- Difference: **513 pcm**

Radial PPF Values

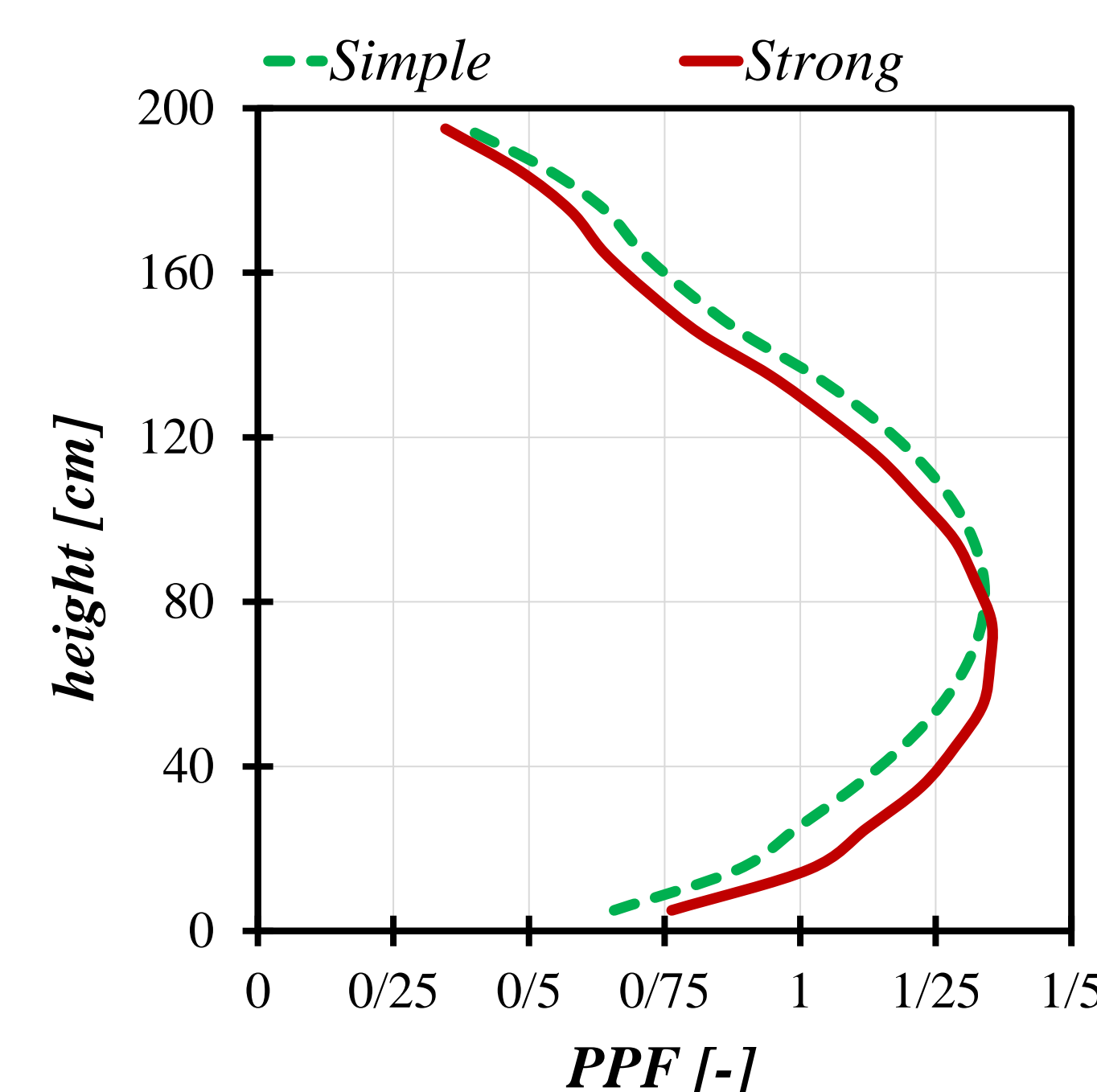
- Simple coupling \rightarrow 1.1303
- Strong coupling \rightarrow 1.1165
- Difference: **1.23%**

Axial PPF Values

- Simple coupling \rightarrow 1.3390
- Strong coupling \rightarrow 1.3530
- Difference: **1.03%**

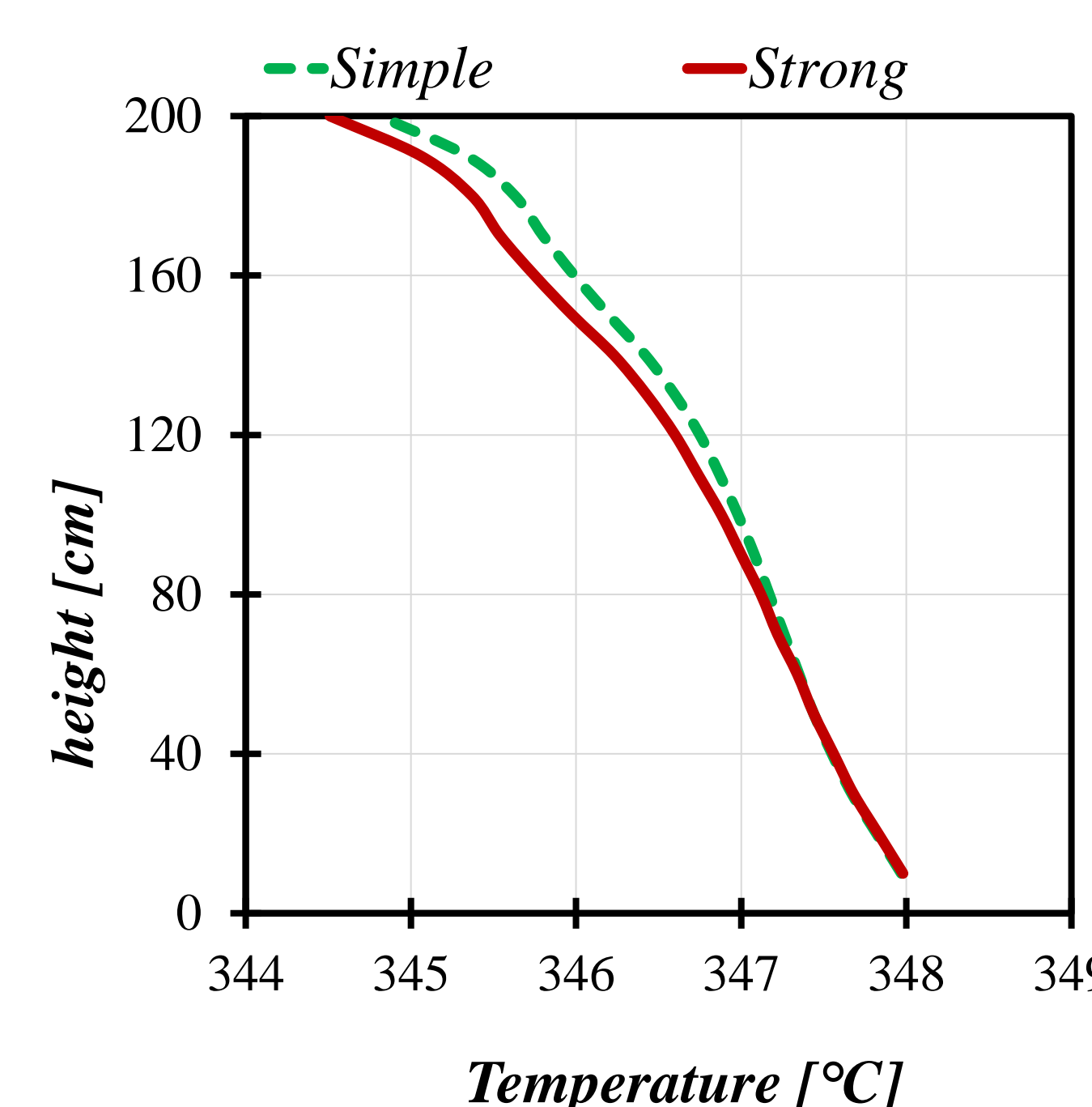
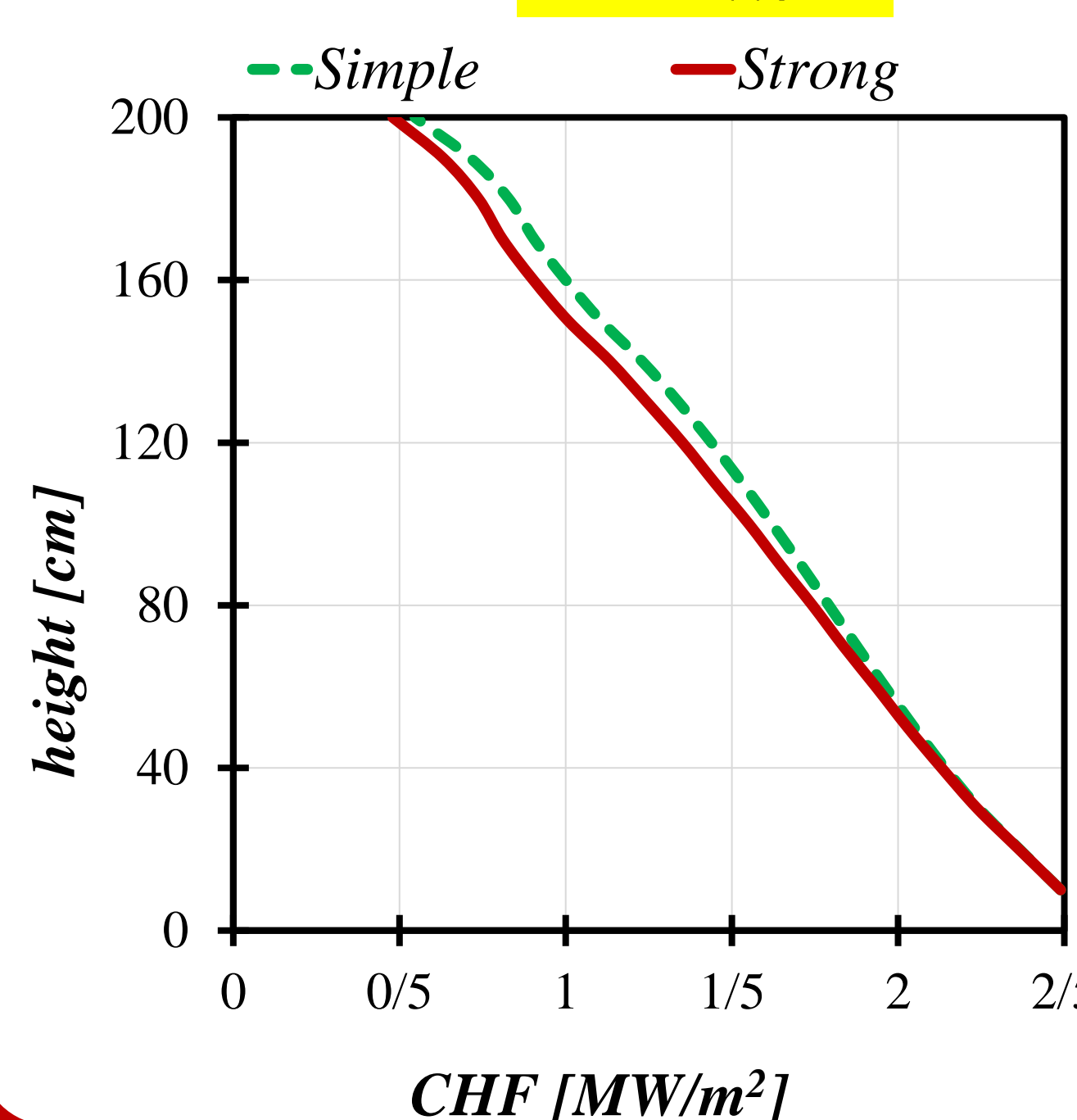
Average CHF Values

- Simple coupling \rightarrow 1.338789 kW/m²
- Strong coupling \rightarrow 1.253989 kW/m²
- Difference: **84.8 kW/m²**



highest CHF temperature difference is around 0.25 K

15.9% relative error in the MDNBR calculations



CONCLUSION

- ✓ The core calculations of a typical **100 MWe** integral SMR
- ✓ **(1)** An **one-way** thermal-hydraulic and neutronic calculation
- ✓ **(2)** Same calculation with **strong iterative coupling** method
- ✓ Neutronic verification: against probabilistic approaches.
- ✓ Thermal-Hydraulic verification: against PARCS code.
- ✓ Both approaches (1) and (2) yielded acceptable average results
- ✓ Data obtained with a simple method are not very trustworthy
- ✓ There is a **15.9%** relative error in the **MDNBR** value

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